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	Filing Date		2005-09-15	
	First Named Inventor	Charles Cantor		
	Art Unit	1634		
	Examiner Name	Stephen Thomas KAPUSHOC		
Attorney Docket Number		701586-053653		

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2	00/50869	WO	A2	2000-08-31	Incyte Pharmaceuticals Inc.		<input type="checkbox"/>
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4	01/75163	WO	A2	2001-10-11	Polygenyx, Inc.		<input type="checkbox"/>
5	01/90399	WO	A2	2001-11-29	Dade Behring Inc.		<input type="checkbox"/>
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	1	BRAUN et al., "Improved Analysis of Microsatellites Using Mass Spectrometry." Genomics 46:18-23, 1997.	<input type="checkbox"/>
	2	BUCHANAN, F. C. et al., "Association of a missense mutation in the bovine leptin gene with carcass fat content and leptin mRNA levels." Genetics Selection Evolution, EDP Sciences, Les Ulis., FR 34(1): 105-116, 2002.	<input type="checkbox"/>
	3	SAUER et al., "Facile method for automated genotyping of single nucleotide polymorphisms by mass spectrometry." Nucleic Acids Research 30(5):1-5, e22, 2002.	<input type="checkbox"/>
	4	Suomalainen, A. and Syvanen, A. C., "Quantitative analysis of human DNA sequences by PCR and solid-phase minisequencing." Mol Biotechnol 15(2):123-131, 2000.	<input type="checkbox"/>

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5	TANG, et al., "Single nucleotide polymorphism analyses by MALDI-TOF MS." International Journal of Mass Spectrometry 226:37-54, 2003.	<input type="checkbox"/>
6	AMEXIS, et al., "Quantitative mutant analysis of viral quasiespecies by chip-based matrix-assisted laser desorption/ionization time-of-flight mass spectrometry." PNAS 98(21):12097-12102, 2001.	<input type="checkbox"/>
7	AMEXIS, et al., "Sequence diversity of Jeryl Lynn strain of mumps virus: quantitative mutant analysis for vaccine quality control." Virology 300(2):171-179, 2002.	<input type="checkbox"/>
8	Becker-André, M. and Hahibrock, K., "Absolute mRNA quantification using the polymerase chain reaction (PCR). A novel approach by a PCR aided transcript titration assay (PATTY)." Nucleic Acids Research 17(22):9437-9446, 1989.	<input type="checkbox"/>
9	BÖCKER, "SNP and mutation discover using base-specific cleavage and MALDI-TOF mass spectrometry." Bioinformatics 19(Suppl 1):i44-i53, 2003.	<input type="checkbox"/>
10	CHENG et al., "Effective amplification of long targets from cloned inserts and human genomic DNA." PNAS 91:5695-5699, 1994.	<input type="checkbox"/>
11	DORIS et al., "Quantitative analysis of gene expression by ion-pair high-performance liquid chromatography." J Chromatography A 806(1):47-60, 1998.	<input type="checkbox"/>
12	ELSO et al., "Mutation Detection Using Mass Spectrometric Separation of Tiny Oligonucleotide Fragments." Genome Research 12:1428-1433, 2002.	<input type="checkbox"/>
13	LANDEGREN et al. "Reading Bits of Genetic Information: Methods for Single-Nucleotide Polymorphism Analysis." Genome Research 8:769-776, 1998.	<input type="checkbox"/>
14	NAKAI et al., "Highly multiplexed genotyping of coronary artery disease-associated SNPs using MALDI-TOF mass spectrometry." Human Mutation 20:133-138, 2002.	<input type="checkbox"/>
15	ROSS et al., "Quantitative Approach to Single-Nucleotide Polymorphism Analysis Using MALDI-TOF Mass Spectrometry." BioTechniques 29(3):620-629, 2000.	<input type="checkbox"/>

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